

FIG. 1

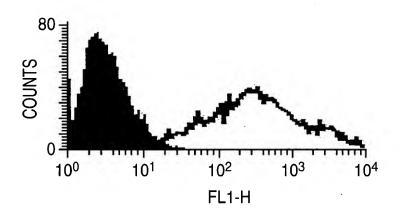


FIG. 2



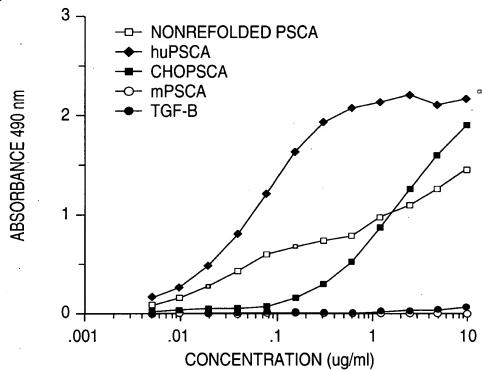


FIG. 3

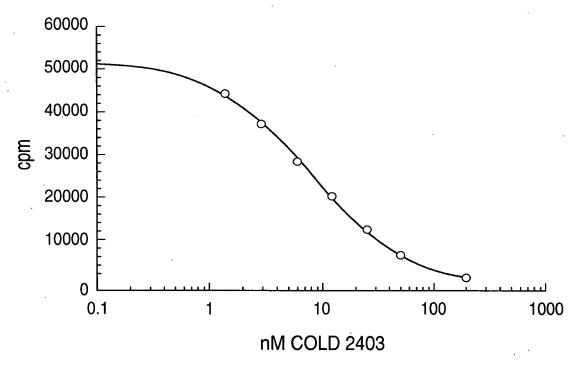


FIG. 4



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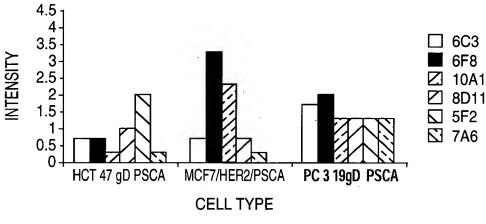
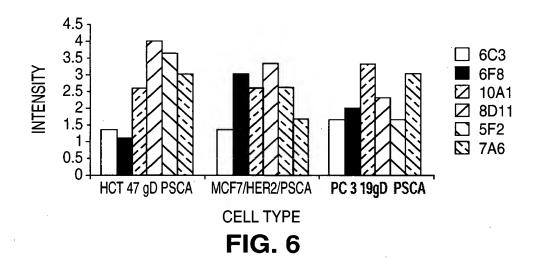


FIG. 5



3.5 3 **SURFACE** 2.5 INTENSITY **INTERNALIZATION** 2 1.5 1 0.5 0 6C3 6F8 10A1 8D11 5F2 7A6 **ANTIBODY** 

**FIG. 7** 

FIG. 8A

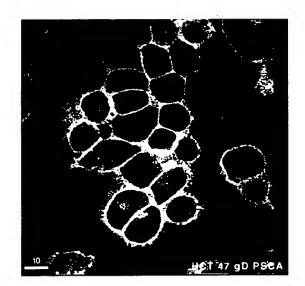


FIG. 8B

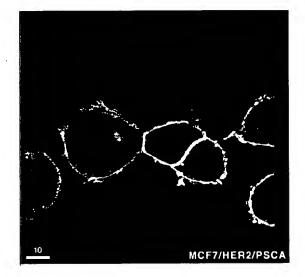


FIG. 8C





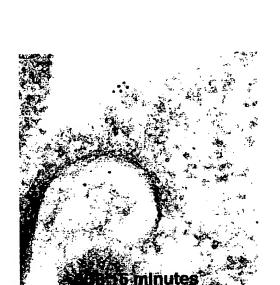


FIG. 9A

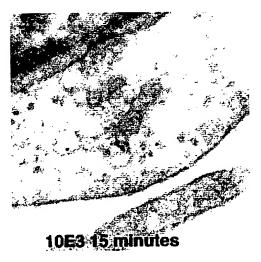


FIG. 9C



FIG. 9B

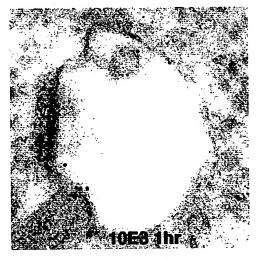
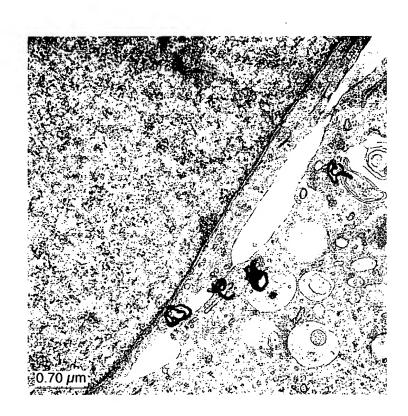


FIG. 9D





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FIG. 10



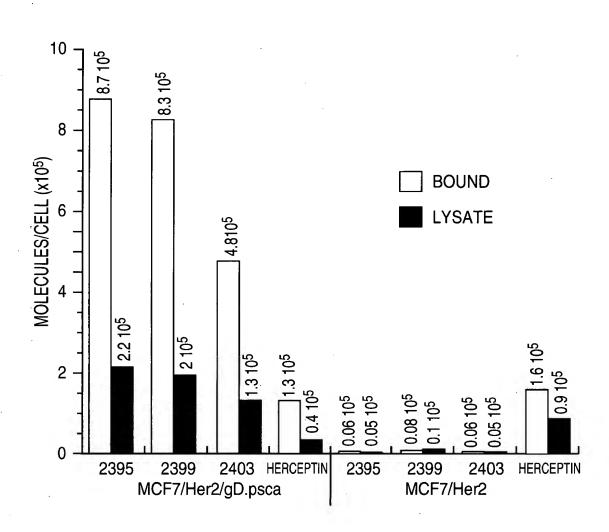


FIG. 11



# HEAVY CHAIN SEQUENCES OF ANTI-PSCA MADS

## CDR1

----ELVKPGAPVKLSCKAS**GYTFTNYWM**WVKORPGRGLEWIG**RIDPSXXXTXXXQTFKD**KA ----PGAELVKPGAPVKLSCKAS**GYTFTNYWMN**WVKQRPGRGLEWIG**RIDPSDSETQYNQTFKD**KA N-QVQVQQPGAELVKPGAPVKLSCKAS**GYTFTNYWLN**WVKQRPGRGLEWIG**RIDPSDSEIHYDQKFKD**KA N-EVQLQQSGPDLEKPGASVKISCKPS**GNSFTGYYIH**WVKQSHGKSLEWIG**RVDPNNGFTSYNQKFKG**KA 2399 2403 2761

### CDR3

ILTVDKSSSTAYMELRSLTSEDSAVYYCVG-NFFDS--WGQGTTLTVSSAKTTGPS-C TLTVDKSSSTAYIQLSSLTSEDSAVYYCAI**TAAIAMDY**WGQGTSVTVSSAKTTGPS-C TLTVDKSSSTAYIQLSSLTSEDSAVYYCAI**TAAIAMDY**WGQGTSVTVSSAKTTGPS-C TLTVDKSSSTAYIQLSSLTSEDSAVYYCAL**TGIYAMAY**WGQGTSVTVSSAKTTGPS-C 2395 2399 2403

# LIGHT CHAIN SEQUENCES OF ANTI-PSCA MAbs

## CDR1

---SVSISCRSSKSLLHSNGNTYLYWFLQRPGQSPQLLIYRMSNLASGVPDRFS N-DIVMTQAAPSVPVTPGESVSISC**RSSKSLLHSNGNTYLY**WFLQRPGQSPQLLIY**RMSNLAS**GVPDRFS N-DVVMTQTPLTLSVTIGQPASISC**KSSQSLLDSDGKTYLN**WLLQRPGQSPKRLIY**LVSTLDS**GVPDRFT 2403 2761 2395

### CDR3

GSGSGTVFTLRISRVEAEDVGVYYC**MQHLESPFT**FGSGTKLEIKR-C GSGSGTAFTLRISRVEAEDVGVYYCLQHLEYPYTFGGGTKLELKR-C 2403

GSGSGTDFTLKISRVEAEDLGVYYCWQGTHFPRTFGGGTKLEIKR-C

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# chimeric 2403 (5F2.4H4.1E3) Light Chain

signal peptide MGWSCIILFLVATATGVHS

VYYCLOHLEYPYTFGGGTKLELK/RTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDS DIVMTQAAPSVPVTPGESVSISCRSSKSLLHSNGNTYLYWFLQRPGQSPQLLIYRMSNLASGVPDRFSGSGSGTAFTLRISRVEAEDVG **IYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC** 

# chimeric 2403 (5F2.4H4.1E3) IgG Heavy Chain

signal peptide MGWSCIILFLVATATGVHS

DVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYT QSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVV LPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYT QVQVQQPGAELVKPGAPVKLSCKASGYTFTNYWLNWVKQRPGRGLEWIGRIDPSDSEIHYDQKFKDKATLTVDKSSSTAYIQLSSLT SEDSAVYYCALTGIYAMAYWGQGTSVTVSSAKTTG/PSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVL

# chimeric 2761 (6B8.1D7.2B3) Fab - Light Chain

DVVMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPGQSPKRLIYLVSTLDSGVPDRFTGSGSGTDFTLKISRVEAEDLGV TVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACET YYCWQGTHFPRTFGGGTKLEIKR [V<sub>1</sub>/Ck junction] HQGLSSPVTKSFNRGEC

# chimeric 2761 (6B8.1D7.2B3) Fab - Heavy chain

EVOLOOSGPDLEKPGASVKISCKPSGNSFTGYYIHWVKQSHGKSLEWIGRVDPNNGFTSYNQKFKGKAILTVDKSSSTAYMELRSLTSE KTTGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSN DSAVYYCVGNFFDSWGQGTTLTVSSA [VH/Cy1 junction] **FKVDKKVEPKSCDKTHT** 

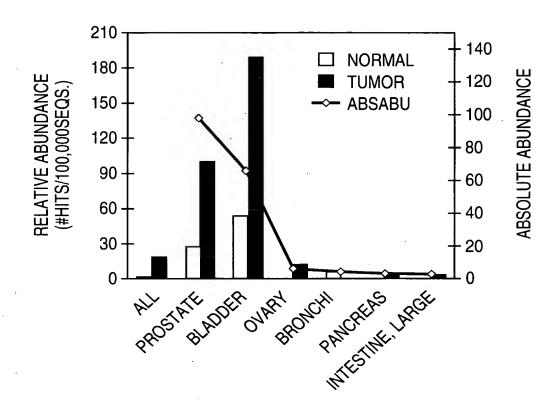
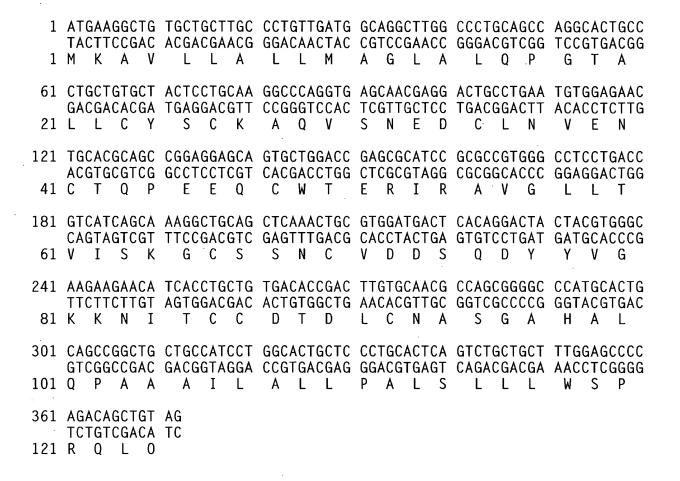


FIG. 14



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**FIG. 15** 

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1 ATGAAGGCTG TGCTGCTTGC CCTGTTGATG GCAGGCTTGG CCCTGCAGCC AGGCACTGCC TACTTCCGAC ACGACGAACG GGACAACTAC CGTCCGAACC GGGACGTCGG TCCGTGACGG 1 M K A V L L A L L M AGLA L 0 P 61 CTGTTGTGCT ACTCCTGCAA GGCCCAGGTG AGCAACGAGG ACTGCCTGAA TGTGGAGAAC GACAACACGA TGAGGACGTT CCGGGTCCAC TCGTTGCTCC TGACGGACTT ACACCTCTTG SNED 21 I I C Y S C K A O VC I N121 TGCACGCAGC CGGAGGAGCA GTGCTGGACC GAGCGCATCC GCGCCGTGGG CCTCCTGACC ACGTGCGTCG GCCTCCTCGT CACGACCTGG CTCGCGTAGG CGCGGCACCC GGAGGACTGG CWT E E QERIR 181 GTCATCAGCA AAGGCTGCAG CTCAAACTGC GTGGATGACT CACAGGACTA CTACGTGGGC CAGTAGTCGT TTCCGACGTC GAGTTTGACG CACCTACTGA GTGTCCTGAT GATGCACCCG S N C V D D S Q D Y

61 V I S K G C S

241 AAGAAGAACA TCACCTGCTG TGACACCGAC TTGTGCAACG CCAGCGGGGC CCATGCCCTG TTCTTCTTGT AGTGGACGAC ACTGTGGCTG AACACGTTGC GGTCGCCCCG GGTACGGGAC T C C D T DLCNA SGA

301 CAGCCAGCTG CTGCCATCCT GGCACTGCTC CCTGCACTCA GCCTGCTGCT TTGGGGCCCC GTCGGTCGAC GACGGTAGGA CCGTGACGAG GGACGTGAGT CGGACGACGA AACCCCGGGG

101 O P A A A I L A L L P A L S LLLWGP

361 AGACAGCTGT AG TCTGTCGACA TC 121 R Q L O

### **FIG. 16**

10 20 30 HUMAN MKAVLLALLMAGLALQPGTALLCYSCKAQVSNEDCLQVENCTQLGEQCWT \*\*\*\*\*\*\*\*\*\*\*\*\* CYNOMOLGUS MKAVLLALLMAGLALQPGTALLCYSCKAQVSNEDCLNVENCTQPEEQCWT 10 20 30

70 90 80 100 ARIRAVGLLTVISKGCSLNCVDDSQDYYVGKKNITCCDTDLCNASGAHAL HUMAN CYNOMOLGUS ERIRAVGLLTVISKGCSSNCVDDSODYYVGKKNITCCDTDLCNASGAHAL 60 70 80 90

110 120 HUMAN OPAAAILALLPALGLLLWGPGOL \*\*\*\*\*\*\*\*\*\* CYNOMOLGUS QPAAAILALLPALSLLLWSPRQL 110 120

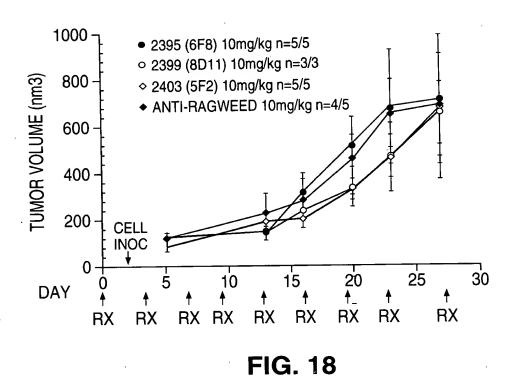
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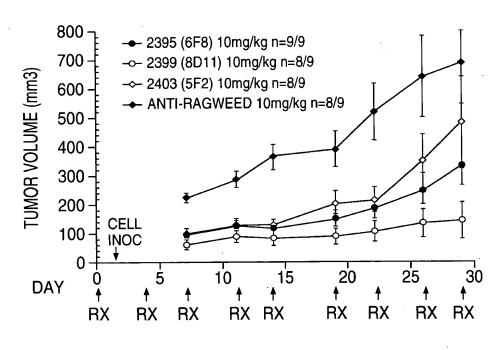


FIG. 19

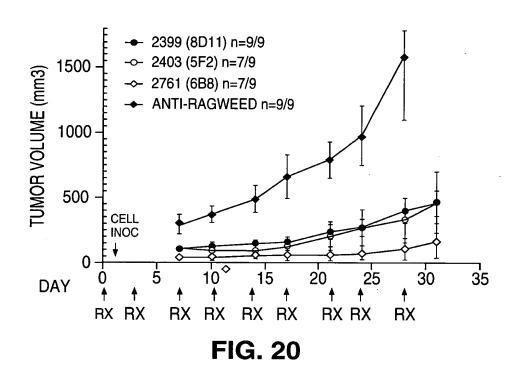
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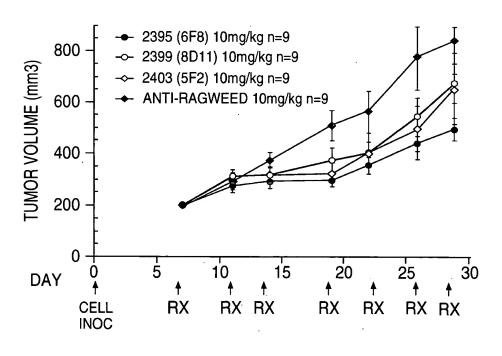


FIG. 21

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FIG. 22

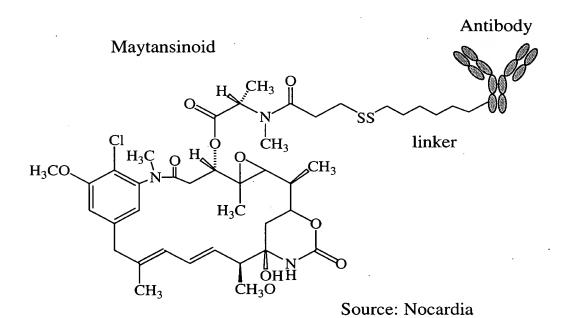


FIG. 23

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Genentech Docket No. P1777R1



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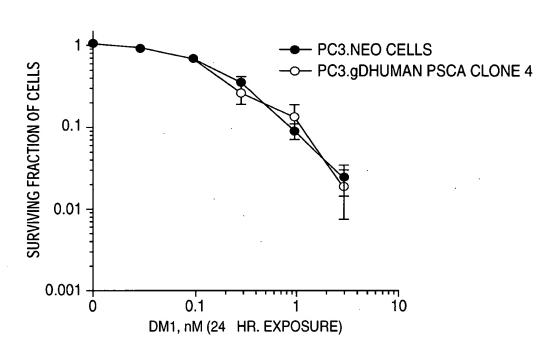


FIG. 24

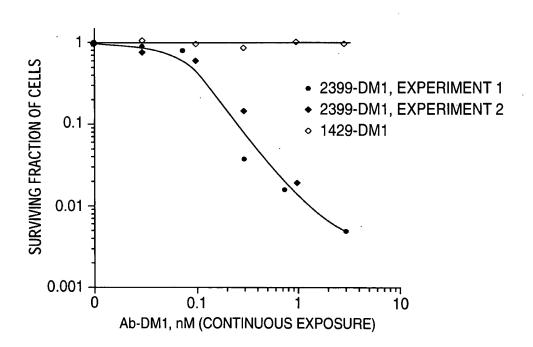


FIG. 25



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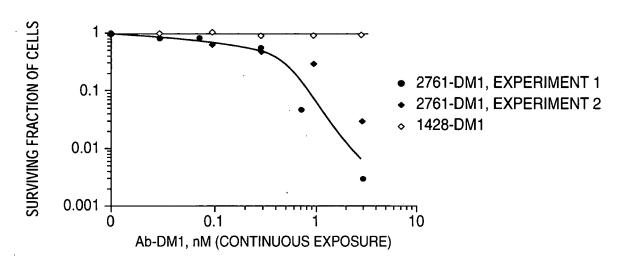


FIG. 26

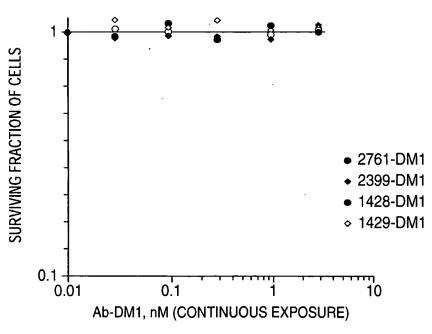


FIG. 27



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- -- 1) 2399 NAKED, 10mg/kg IP, 2X WEEK, n=7/7
- 2) 6B8 NAKED, 10mg/kg IP, 2X WEEK, n=7/7
- → 3) ANTI-RAGWEED, 10mg/kg IP, 2X WEEK, n=7/7
- → 4) 2399-DM1, 75 ug/kg IV, 2X WEEK, n=3/7
- → 5) 6B8-DM1, 75 ug/kg IV, 2X WEEK, n=3/7
- -□- 6) ANTI-RAGWEED-DM1, 75 ug/kg IV, 2X WEEK, n=6/7

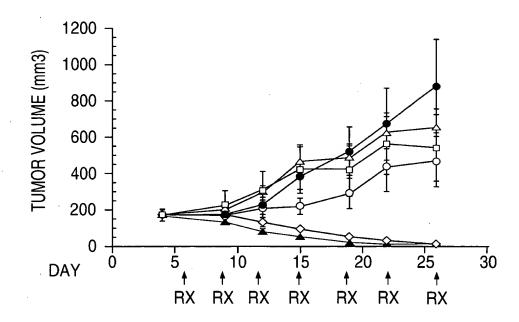


FIG. 28